
Sequence Listing was accepted.

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Reviewer: markspencer

Timestamp: [year=2011; month=12; day=21; hr=11; min=0; sec=3; ms=909;]

Validated By CRFValidator v 1.0.3

Application No: 10582115 Version No: 2.0

Input Set:

Output Set:

Started: 2011-12-14 16:42:42.173

Finished: 2011-12-14 16:42:45.679

Elapsed: 0 hr(s) 0 min(s) 3 sec(s) 506 ms

Total Warnings: 46

Total Errors: 0

No. of SeqIDs Defined: 118

Actual SeqID Count: 118

Error code		Error Description
W	213	Artificial or Unknown found in <213> in SEQ ID (1)
W	213	Artificial or Unknown found in <213> in SEQ ID (2)
W	213	Artificial or Unknown found in <213> in SEQ ID (3)
W	213	Artificial or Unknown found in <213> in SEQ ID (4)
W	213	Artificial or Unknown found in <213> in SEQ ID (5)
W	213	Artificial or Unknown found in <213> in SEQ ID (6)
W	213	Artificial or Unknown found in <213> in SEQ ID (7)
W	213	Artificial or Unknown found in <213> in SEQ ID (8)
W	213	Artificial or Unknown found in <213> in SEQ ID (9)
W	213	Artificial or Unknown found in <213> in SEQ ID (10)
W	213	Artificial or Unknown found in <213> in SEQ ID (11)
W	213	Artificial or Unknown found in <213> in SEQ ID (12)
W	213	Artificial or Unknown found in <213> in SEQ ID (13)
W	213	Artificial or Unknown found in <213> in SEQ ID (14)
W	402	Undefined organism found in <213> in SEQ ID (20)
W	402	Undefined organism found in <213> in SEQ ID (21)
W	402	Undefined organism found in <213> in SEQ ID (22)
W	402	Undefined organism found in <213> in SEQ ID (23)
W	402	Undefined organism found in <213> in SEQ ID (24)
W	402	Undefined organism found in <213> in SEQ ID (40)

Input Set:

Output Set:

Started: 2011-12-14 16:42:42.173

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Total Warnings: 46

Total Errors: 0

No. of SeqIDs Defined: 118

Actual SeqID Count: 118

Error code		Error Description
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W	402	Undefined organism found in <213> in SEQ ID (42)
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M	402	Undefined organism found in <213> in SEQ ID (46)
W	402	Undefined organism found in <213> in SEQ ID (47)
W	402	Undefined organism found in <213> in SEQ ID (48)
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W	402	Undefined organism found in <213> in SEQ ID (57)
W	402	Undefined organism found in <213> in SEQ ID (58)
W	402	Undefined organism found in <213> in SEQ ID (59) This error has occured more than 20 times, will not be displayed
W	213	Artificial or Unknown found in <213> in SEQ ID (118)

SEQUENCE LISTING

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<110> ZOLLO, MASSIMO
<120> USE OF ENZYMATIC INHIBITORS OF H-PRUNE FOR THE PREVENTION AND
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<130> 026073-00006
<140> 10582115
<141> 2011-12-14
<150> PCT/IT2004/000689
<151> 2004-12-10
<150> IT RM2003A000572
<151> 2003-12-11
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<210> 8
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<213> Artificial Sequence
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33

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<210> 10 <211> 29 <212> PRT <213> Artificial Sequence <220>

<223> Description of Artificial Sequence: Synthetic permeable inhibitor of h-prune H1 sequence

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20 25

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<211> 15

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<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic
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                                    10
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<400> 12

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Ala Thr Asp Thr Gly Ser Phe Arg Trp Ala Ser Val
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<213> Mycobacterium tuberculosis
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Thr Val Asn Leu Ala Ala Val Ala Ser Gly Phe Gly Gly Gly His
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Asp Leu Ile Leu Cys His Gln Thr Ala Asp Phe Asp Val Leu Gly Ala
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25 20

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Lys Ala Ala
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Arg Gln Val Ala Ile Tyr Asp His His Leu Asn Ser Pro
1
                5
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                                                         15
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                                    10
Asp Ile Ser Leu Ser Met Val Glu Ala Ser Val Met Ala Leu Gly Ile
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                                25
                                                     30
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<212> PRT
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                                                         15
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Ala Gln Ala Ala Val Asn Leu Arg Asp Val 20 25

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                                     10
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Gln Gln Gly Leu Phe His Leu Ile Lys Ala Asn Phe
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                                 25
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Glu Ala Leu Ala Ile Val Val Asp Ala Asn Tyr Lys Asn Arg Ile Glu
1
                                     10
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Leu Arg Glu
<210> 27
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<400> 27
Lys Ala Val Leu Arg Ile Asp His His Pro Asn Glu Asp
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<211> 44
<212> PRT
<213> Mycoplasma genitalium
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Ser Tyr Val Ala Cys Cys Glu Gln Ile Val Glu Met Ala Thr Val Ala
                                     10
                                                         15
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Lys Trp Thr Ile Pro Pro Val Ala Ala Thr Leu Leu Tyr Ile Gly Ile
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                                                     30
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Tyr Thr Asp Ser Asn Arg Phe Leu Tyr Ser Asn Thr

40

35

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<212> PRT
<213> Bacillus subtilis
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                                    10
                                                        15
1
Gln Cys Gly Leu Thr Glu Ile Leu Arg Glu Thr Tyr
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                                25
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Asp Gln Arg
<210> 32
<211> 13
<212> PRT
<213> Bacillus subtilis
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Ala Lys Leu Met Lys Ile Asp His His Pro Asn Glu Asp
                5
1
                                    10
<210> 33
<211> 44
<212> PRT
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<213> Bacillus subtilis

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10

15

Val Gly Asp Thr Gly Arg Phe Leu Phe Pro Asn Thr 35

5

<210> 34

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<212> PRT

<213> Bacillus subtilis

<400> 34

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1 10 15

Pro Leu Ala Ser Gly Ala Ser Ile Tyr Ser Trp
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<210> 35

<211> 28

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<213> Archaeoglobus fulgidus

<400> 35

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Ala Tyr Ala Leu Arg Glu Ile Ala Lys Gln Phe Asp 20 25

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<213> Archaeoglobus fulgidus

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Ser Ile Pro

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                                    10
                                                         15
1
Lys Ile Thr Pro Ser Lys Ile Leu Ala Thr Ala Leu Phe Phe Gly Ile
            20
                                25
                                                     30
Lys Ser Glu Thr Asp Glu Phe Lys Arg Asn Thr Arg
        35
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<213> Archaeoglobus fulgidus
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1
                                    10
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            20
                                25
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<213> Methanocaldococcus jannaschii
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1
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                                                         15
Arg Ala Ile Leu Gln Lys Leu Ala Glu Arg Leu Asn
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                                25
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                                                        15
1
Ile Lys Glu
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Asp Lys Ile Ile Leu Asp His His Gln Pro Glu Glu
                                    10
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<213> Methanocaldococcus jannaschii
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1
                                                        15
                                    10
Ile Asn Asn Asp Trp Ile Asp Leu Ala Lys Tyr Ala Val Leu Gly Ala
            20
                                25
                                                     30
Val Gly Asp Ile Gln Asn Ile Glu Gly Lys Leu Ile
        35
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<213> Methanocaldococcus jannaschii
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                                                        15
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            20
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<211> 28

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                                                         15
1
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            20
                                25
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1
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Pro Ala Ile
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<213> Methanocaldococcus jannaschii
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                                    10
                                                         15
1
Met Asp Phe Glu Ala Phe Tyr Leu Arg Phe Met Asp Gly Lys Gly Ile
            20
                                25
                                                     30
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<213> Methanocaldococcus jannaschii

<213> Methanocaldococcus jannaschii

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1
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            20
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1
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Glu Tyr Leu
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1
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